

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/549,352
Source: PET
Date Processed by STIC: 09/28/2005

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER** **VERSION 4.2.2 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

10/549,352

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing, the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



PCT

RAW SEQUENCE LISTING

DATE: 09/28/2005

PATENT APPLICATION: US/10/549,352

TIME: 13:53:52

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Output Set: N:\CRF4\09282005\J549352.raw

6 <110> APPLICANT: DRAKE, Caroline Rachel
 7 PAINE, Jacqueline Ann Mary
 8 SHIPTON, Catherine Ann
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 C--> 16 <140> CURRENT APPLICATION NUMBER: US/10/549,352
 C--> 16 <141> CURRENT FILING DATE: 2005-09-14
 16 <150> PRIOR APPLICATION NUMBER: PCT/GB2004/001241
 17 <151> PRIOR FILING DATE: 2004-03-22
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Does Not Comply
 Corrected Diskette Needed
 Cpg-1, 3, 5,

Invalid Response,
 Organism can be either Artificial,
 Unknown or Genus species, pls
 see gloss #10 on Error Sum-
 mary sheet.

RAW SEQUENCE LISTING

DATE: 09/28/2005

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Output Set: N:\CRF4\09282005\J549352.raw

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Same Error

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Input Set : A:\70237USPCT SEQUENCE LISTING.txt

Output Set: N:\CRF4\09282005\J549352.raw

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340	aacaccaaaa	tatgcacaag	gcagtttggt	gtattctgta	gtacagacaa	aactaaaagt	3120
342	aatgaaagaa	gatgtggtgt	tgaaaaagga	aacaatatca	tgagtaatgt	gtgagcatta	3180
344	tgggaccacg	aaataaaaag	aacattttga	tgagtcgtgt	atcctcgatg	agcctcaaaa	3240
346	gttctctcac	cccggataag	aaacccttaa	gcaatgtgca	aagtgtgcat	tctccactga	3300
348	cataatgcaa	aataagatat	catcgatgac	atagcaactc	atgcataata	tcatgcctct	3360
350	ctcaacctat	tcattctctac	tcattctacat	aagtatcttc	agctaaatgt	tagaacataa	3420
352	accataaagt	cacgtttgat	gagtattagg	cgtgacacat	gacaaatcac	agactcaagc	3480
354	aagataaagc	aaaatgatgt	gtacataaaa	ctccagagct	atatgtcata	ttgcaaaaag	3540
356	aggagagcct	ataagacaag	gcatgactca	caaaaattca	tttgcccttc	gtgtcaaaaa	3600
358	gaggagggct	ttacattatc	catgtcatat	tgcaaaagaa	agagagaaag	aacaacacaa	3660
360	tgctgcgtca	attatacata	tctgtatgtc	catcattatt	catccacctt	tcgtgtacca	3720
362	cacttcatat	atcatgagtc	acttcatgtc	tggacattaa	caaactctat	cttaacattt	3780
364	agatgcaaga	gcctttatct	cactataaet	gcacgatgat	ttctcattgt	ttctcacaaa	3840
366	aagcattcag	ttcattagtc	ctacaacaac	gaattcggct	tcccgggtac	agggtaaatt	3900
368	tctagttttt	ctccttcatt	ttcttggtta	ggaccttttt	ctctttttat	ttttttgagc	3960
370	tttgatcttt	ctttaaactg	atctattttt	taattgattg	gttatcgtgt	aaatattaca	4020
372	tagctttaac	tgataatctg	attactttat	ttcgtgtgtc	tttgatcatc	ttgatagtta	4080
374	cagaaccgct	gactctagag	aagccattta	aatcgccgce	accatggcca	tcatactcgt	4140
376	acgagcagcg	tcgcccgggc	tctccgcccgc	cgacagcacc	agccaccagg	ggactctcca	4200
378	gtgctccacc	ctgctcaaga	cgaagaggcc	ggcgccgccc	cgggtggatgc	cctgctcgct	4260
380	ccttggcctc	cacccgtggg	aggctggccg	tccctcccc	gccgtctact	ccagcctcgc	4320
382	cgtcaacccg	gcgggagagg	cgtcgtctc	gtccgagcag	aagggtctacg	acgtcgtgct	4380

RAW SEQUENCE LISTING

DATE: 09/28/2005

PATENT APPLICATION: US/10/549,352

TIME: 13:53:52

Input Set : A:\70237USPCT SEQUENCE LISTING.txt

Output Set: N:\CRF4\09282005\J549352.raw

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384 caagcagggc gcattgctca aacgccagct gcgcacggcg gtcctcgacg ccaggcccca 4440
386 ggacatggac atgccacgca acgggctcaa ggaagcctac gaccgctgcg gcgagatctg 4500
388 tgaggagtat gccaagacgt tttacctcgg aactatgttg atgacagagg agcggcgccg 4560
390 cgccatatgg gccatctatg tgtggtgtag gaggacagat gagctttag atgggcaaaa 4620
392 cgccaactac attacaccaa cagctttgga ccggtgggag aagagacttg aggatctgtt 4680
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396 catagacatt cagccattca gggacatgat tgaagggatg aggagtgtac ttaggaagac 4800
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404 tggagaggat gctagacgag gaaggatata tttaccacaa gatgagcttg cacaggcagg 5040
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408 gaggcagatc aagagggcca ggatgttttt tgaggaggca gagagagggg taactgagct 5160
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418 agtttcttaa gattgaatcc tgttgccggg cttgcgatga ttatcatata atttctgttg 5460
420 aattacgta agcatgtaat aattaacatg taatgcatac cgttatttat gagatgggtt 5520
422 tttatgatta gagtcccgca attatacatt taatacgcga tagaaaacaa aatatagcgc 5580
424 gcaaaactag ataaattatc gcgcgcgggtg tcactatgtt tactagatcg 5630
427 <210> SEQ ID NO: 3
429 <211> LENGTH: 5180
431 <212> TYPE: DNA
433 <213> ORGANISM: SYNTHETIC - 12422
437 <400> SEQUENCE: 3
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440 tattctgtag tacagacaaa actaaaagta atgaaagaag atgtggtgtt agaaaaggaa 120
442 acaatatcat gagtaatgtg tgagcattat gggaccacga aataaaaaga acattttgat 180
444 gagtcgtgta tctctgatga gctcaaaag ttctctcacc ccgataaga aacccttaag 240
446 caatgtgcaa agtttgcat ctccactgac ataatagcaa ataatagatc atcgatgaca 300
448 tagcaactca tgcattcat catgcctctc tcaacctatt cattctact catctacata 360
450 agtatcttca gctaaatgtt agaacataaa cccataagtc acgtttgat agtattaggc 420
452 gtgacacatg acaaatcaca gactcaagca agataaagca aatgatgtg tacataaaac 480
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456 aaaaattcat ttgcctttcg tgtcaaaaag aggagggtt tacattatcc atgtcatatt 600
458 gcaaaagaaa gagagaaaga acaacacaat gctgcgtcaa ttatacatat ctgtatgtcc 660
460 atcattattc atccaccttt cgtgtaccac acttcatata tcatgagtc cttcatgtct 720
462 ggacattaac aaactctatc ttaacattta gatgcaagag cctttatctc actataaatg 780
464 caccgatgatt tctcattgtt tctcacaana agcattcagt tcattagtc tacaacaacg 840
466 aattcggctt cccaaatcgc cgccaccatg gcttctatga tctctcttc cgctgtgaca 900
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470 tccatgactg gattcccagt gaagaaggte aacactgaca ttacttccat tacaagcaat 1020
472 ggtggaagag taaagtgcac gaaaccaact acggttaatt gtgcaggctt cgggtggcctg 1080
474 gcaactggcaa ttcgtctaca agctgcgggg atccccgtct tactgttgga acaacgtgat 1140
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478 acggttatca ccgatcccag tgccattgaa gaactgtttg cactggcagg aaaacagtta 1260
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482 aaggtcttta attacgataa cgatcaaacc cggctcgaag cgcagattca gcagttaat 1380

```

*Same Error
Organism Can be either
Artificial,
Unknown,
or Genus
Species.
Pls see
Item #
ID on
Error Sum-
mary sheet.*

*The type of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.*

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/28/2005
PATENT APPLICATION: US/10/549,352 TIME: 13:53:53

Input Set : A:\70237USPCT SEQUENCE LISTING.txt
Output Set: N:\CRF4\09282005\J549352.raw

Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"
or "Unknown". Please explain source of genetic material in <220> to <223>
section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)
(Sec.1.823 of new Rules)

Seq#:22

VERIFICATION SUMMARY

DATE: 09/28/2005

PATENT APPLICATION: US/10/549,352

TIME: 13:53:53

Input Set : A:\70237USPCT SEQUENCE LISTING.txt

Output Set: N:\CRF4\09282005\J549352.raw

L:16 M:270 C: Current Application Number differs, Replaced Current Application No
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2513 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
L:2517 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:22, <213>
ORGANISM:Artificial Sequence
L:2517 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:22, <213>
ORGANISM:Artificial Sequence
L:2517 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:22,Line#:2517